

## Aberystwyth University

### Erratum to 'Combining bioinformatics, cheminformatics, functional genomics and whole organism approaches for identifying epigenetic drug targets in *Schistosoma mansoni*' [IJP Drugs Drug Resist. 8 (2018) 559–570]

Padalino, Gilda; Ferla, Salvatore; Brancale, Andrea; Chalmers, Iain; Hoffmann, Karl

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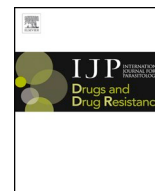
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## Erratum to ‘Combining bioinformatics, cheminformatics, functional genomics and whole organism approaches for identifying epigenetic drug targets in *Schistosoma mansoni*’ [IJP Drugs Drug Resist. 8 (2018) 559–570]



Gilda Padalino<sup>a</sup>, Salvatore Ferla<sup>b</sup>, Andrea Brancale<sup>b</sup>, Iain W. Chalmers<sup>a</sup>, Karl F. Hoffmann<sup>a,\*</sup>

<sup>a</sup> The Institute of Biological, Environmental and Rural Sciences (IBERS), Aberystwyth University, SY23 3DA, Wales, UK

<sup>b</sup> School of Pharmacy and Pharmaceutical Sciences, Cardiff University, Cardiff, CF10 3NB, United Kingdom

The publisher regrets to inform that there was errors in the column headers of table 1 published in the original article. The corrected table 1 is as below:

Table 1  
L8 and L10 antischistosomal activity and cytotoxicity summary.

Compound	EC <sub>50</sub> on schistosomula phenotype	EC <sub>50</sub> on schistosomula motility	EC <sub>50</sub> on juveniles	EC <sub>50</sub> on adult worms	CC <sub>50</sub> on HepG2 cells	SI CC <sub>50</sub> /EC <sub>50</sub> schistosomula (phenotype/motility)	SI CC <sub>50</sub> /EC <sub>50</sub> juveniles	SI CC <sub>50</sub> /EC <sub>50</sub> adult worms
<b>L8</b>	4.19 μM	5.86 μM	3.91 μM	> 50 μM	17.79 μM	4.25/3.04	4.55	< 0.36
<b>L10</b>	2.49 μM	1.46 μM	2.28 μM	14.18 μM	21.05 μM	8.45/14.42	9.23	1.48

The publisher would like to apologise for any inconvenience caused.

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\* Corresponding author.

E-mail addresses: [gip3@aber.ac.uk](mailto:gip3@aber.ac.uk) (G. Padalino), [ferlaS1@cardiff.ac.uk](mailto:ferlaS1@cardiff.ac.uk) (S. Ferla), [brancaleA@cardiff.ac.uk](mailto:brancaleA@cardiff.ac.uk) (A. Brancale), [iwc@aber.ac.uk](mailto:iwc@aber.ac.uk) (I.W. Chalmers), [krh@aber.ac.uk](mailto:krh@aber.ac.uk) (K.F. Hoffmann).

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